

B18 mRNA (arbitrary units)

FIG. 1

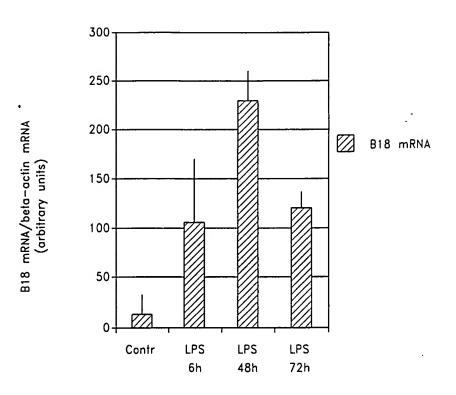


FIG. 2

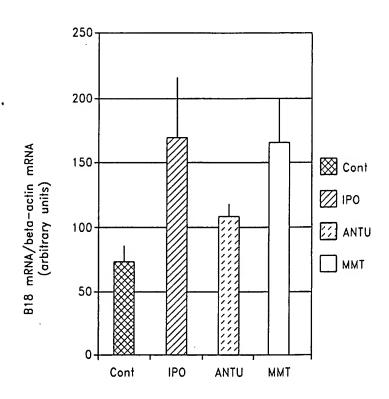


FIG. 3



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#### Gene (chromosome 11q12-13)

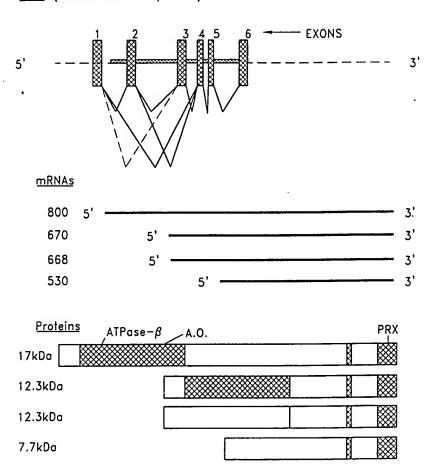


FIG. 4

CLUSTAL V alignment of human and rat B18 amino acid sequences (Identity: 908, Homology: 97.5%):
B18hum MAPIKVGDAIPAVEVFEGEPGNKVNLAELFKGKKGVLFGVPGAFTPGCSK = SEQIDNO1 B18rat MAPIKVGDTIPSVEVFEGEPGKKVNLAELFKDKKGVLFGVPGAFTPGCSK ************************************
B18hum THLPGFVEQAEALKAKGVQVVACLSVNDAFVTGEWGRAHKAEGKVRLLAD B18rat THLPGFVEQAGALKAKGAQVVACLSVNDVFVTAEWGRAHQAEGKVQLLAD ********* ****** ********************
B18hum PTGAFGKETDLLLDDSLVSIFGNRRLKRFSMVVQDGIVKALNVEPDGTGL B18rat PTGAFGKETDLLLDDSLVSLFGNRRLKRFSMVIDKGVVKALNVEPDGTGL ***********************************
B18hum TCSLAPNIISQL B18rat TCSLAPNILSQL ********
CLUSTAL V alignment of human and mouse B18 amino acid sequences (Identity: 91%, Homology: 96%)
B18hum MAPIKVGDAIPAVEVFEGEPGNKVNLAELFKGKKGVLFGVPGAFTPGCSK MAPIKVGDAIPSVEVFEGEPGKKVNLAELFKGKKGVLFGVPGAFTPGCSK ************************************
B18hum THLPGFVEQAEALKAKGVQVVACLSVNDAFVTGEWGRAHKAEGKVRLLAD THLPGFVEQAGALKAKGAQVVACLSVNDVFVIEEWGRAHQAEGKVRLLAD
B18hum PTGAFGKETDLLLDDSLVSIFGNRRLKRFSMVVQDGIVKALNVEPDGTGL B18mouse PTGAFGKATDLLLDDSLVSLFGNRRLKRFSMVIDNGIVKALNVEPDGTGL
B18hum TCSLAPNIISQL B18mouse TCSLAPNILSQL ********
CLUSTAL V alignment of human and rat cDNA sequences (identity: 612/780,78.5%):
B18hum GCCAGGAGGCGGAGTGGAAGTGGCCGTGGGGCGGGTATGGGACTAGCTGGTGCGTCCTAGGCAG , ** *** ***
B18hum CGTGTGCGCCCTGAGACGCTCAGCGGGCTATATACTCGTCGGTGGGGCCG B18rat CATAGCCGGATCGGTGCTCCGTGCATCGGCTACTTGGAC- * * * * * * * * * * * * * * * * * * *
B18hum GCGGTCAGTCTGCGGCAGCGGCAGCAGACGGTGCAGTGAAGGAGAGTGG B18ratGTGCGTGGCAGGCAGGCAGGCCGGAAAGGAGCAGGTTGG ** * **** * *** * * * * * * * * * * *

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	GCGTCTGGCGGGGTCCGCAGTTTCAGCAGAGCCGCTGCAGCCATGGCCCC GAGTGTGGTGGGGCCCGCAGCTTCAGCAGTGCCGCGGTGACTATGGCCCC * ** *** *** **** ***** ***** * * *
B18hum B18rat	AATCAAGGTGGGAGATGCCATCCCAGCAGTGGAGGTGTTTGAAGGGGAGCGATCAAGGTGGAGACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAACCATCAGACACACAC
	CAGGGAACAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAGGGTGTGCTGGGAAAGAAGGTGAACTTGGCAGAGCTGTTCAAGGACAAGAAAGGTGTT
B18hum B18rat	CTGTTTGGAGTTCCTGGGGCCTTCACCCCTGGATGTTCCAAGACACACCT TTGTTTGGAGTCCCTGGGCATTTACACCTGGCTGTTCCAAGACCCATCT ********* ******* ** ****** **********
	GCCAGGGTTTGTGGAGCAGGCTGAGGCTCTGAAGGCCAAGGGAGTCCAGG GCCTGGGTTTGTGGAGCAAGCCGGAGCTCTGAAGGCCAAGGGAGCACAAG *** ********* ** * * ******** ** * * *
B18hum B18rat	TGGTGGCCTGTCTGAGTGTTAATGATGCCTTTGTGACTGGCGAGTGGGGC TGGTGGCCTGTCTGAGTGTTAATGATGTCTTCGTGACTGCAGAGTGGGGT *****************************
B18hum B18rat	CGAGCCCACAAGGCGGAAGGCAAGGTTCGGCTCCTGGCTGATCCCACTGG CGAGCCCACCAGGCAGAAGGCAAGGTTCAGCTCCTGGCTGACCCCACTGG
B18hum B18rat	GGCCTTTGGGAAGGAGACAGACTTATTACTAGATGATTCGCTGGTGTCCA AGCTTTTGGAAAGGAGACAGATTTACTACTAGATGATTCTTTTGGTGTCTC ** **** ***************************
B18hum B18rat	TCTTTGGGAATCGACGTCTCAAGAGGTTCTCCATGGTGGTACAGGATGGC TCTTTGGGAATCGTCGGCTAAAAAGGTTCTCCATGGTGATAGACAAGGGC *****************************
	ATAGTGAAGGCCCTGAATGTGGAACCAGATGGCACAGGCCTCACCTGCAG GTAGTAAAGGCACTGAACGTGGAGCCGGATGGCACAGGCCTCACCTGCAG **** **** ***** *********************
B18hum B18rat	CCTGGCACCCAATATCATCTCACAGCTCTGAGGCCCTGGGCCAGATTACT CCTGGCCCCCAACATCCTCTCACAACTCTGAGGCCCTGA-CCAGAATG ***** **** *** *** **** ******* ***** ****
	TCCTCCACCCTCCTATCTCACCTGCCCAGCCCTGTGCTGG-GGCCCTG TCCTCTGACTCTCCC-ATCTCCTCCACCCAGCTCTGGGCCAAAGGCCCAG *****
B18hum	CATTGGCCAGATTTCTGC TACCTCCTTACCTGAGGGCCACTGGAATGGAA
-	AATAAACACTTGTGGTTTGCGGAAAAAAAAAAAAAAAAA

FIG. 5B

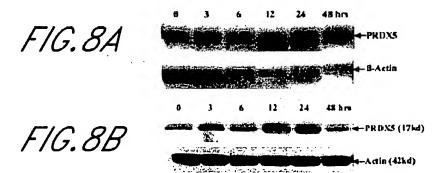
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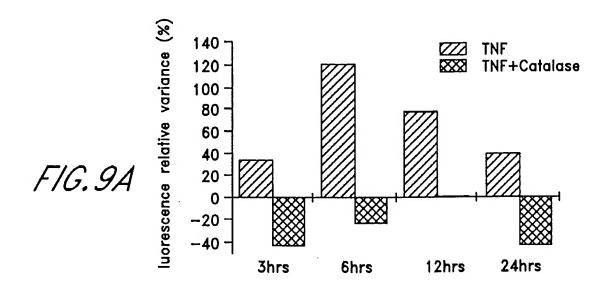
CLUSTAL V alignment of human and mouse cDNA sequences (identity:552/675, 81.8%)
B18hum GCCAGGAGGCGGAGTGGAAGTGGCCGTGGGGCGGGTATGGGACTAGCTGG B18mouse
B18hum CGTGTGCGCCCTGAGACGCTCAGCGGGCTATATACTCGTCGGTGGGGCCG B18mouseTGCTCCGTGCATCGACGTGCTTG
B18hum GCGGTCAGTCTGCGGCAGCGGCAGCAAGACGGTGCAGTGAAGGAGAGGTGG B18mouse GCAGGCAGAGCAGGCCGGAAAGAAGCAGGTTGG ** * * * * * * * * * * * * * * * * *
B18hum GCGTCTGGCGGGGTCCGCAGTTTCAGCAGAGCCGCTGCAGCCATGGCCCC B18mouse GAGTGTGGCGGAGCCCGCAGCTTCAGCAGCTCCGCGGTGACCATGGCCCC  * * * * * * * * * * * * * * * * *
B18hum AATCAAGGTGGGAGATGCCATCCCAGCAGTGGAGGTGTTTGAAGGGGAGC B18mouse GATCAAGGTGGAGATGCCATTCCCTCAGTGGAGGTATTTGAAGGGGAAC *****************************
B18hum CAGGGAACAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAGGGTGTG B18mouse CGGGAAAGAAGGTGAACTTGGCAGAGCTGTTCAAGGGCAAGAAAGGTGTT  * ** ** ******** ***************
B18hum CTGTTTGGAGTTCCTGGGGCCTTCACCCCTGGATGTTCCAAGACACACCT B18mouse TTGTTTGGAGTCCCTGGGGCATTTACACCTGGCTGTTCTAAGACCCACCT ********* ****** ** ** ** ***** ***** ****
B18hum GCCAGGGTTTGTGGAGCAGGCTGAGGCTCTGAAGGCCAAGGGAGTCCAGG B18mouse GCCTGGGTTTGTGGAGCAAGCTGGAGCTCTGAAGGCTAAGGGAGCGCAGG *** ********** **** ******* ****** *****
B18hum TGGTGGCCTGTCTGAGTGTTAATGATGCCTTTGTGACTGGCGAGTGGGGC B18mouse TGGTGGCCTGTCTGAGCGTTAATGACGTCTTTGTGATTGAAGAGTGGGGT ******************
B18hum CGAGCCCACAAGGCGGAAGGCAAGGTTCGGCTCCTGGCTGATCCCACTGG B18mouse CGAGCCCACCAGGCAGAAGGCAAGGTTCGGCTCCTGGCTGACCCCACTGG
B18hum GGCCTTTGGGAAGGAGACAGACTTATTACTAGATGATTCGCTGGTGTCCA B18mouse AGCCTTTGGGAAGGCGACAGACTTATTATTGGATGATTCTTTGGTGTCTC *************************
B18hum TCTTTGGGAATCGACGTCTCAAGAGGTTCTCCATGGTGGTACAGGATGGC B18mouse TCTTTGGGAATCGTCGGCTGAAAAGGTTCTCCATGGTGATAGACAACGGC
B18hum ATAGTGAAGGCCCTGAATGTGGAACCAGATGGCACAGGCCTCACCTGCAG B18mouse ATAGTGAAGGCACTGAACGTGGAGCCAGATGGCACAGGCCTCACCTGCAG ***********************************
B18hum CCTGGCACCCAATATCATCTCACAGCTCTGAGGCCCTGGGCCAGATTACT B18mouse CCTGGCCCCCAACATCCTCTCCCAACTCTGAGGCCCTGG-CCAGATG ***** **** **** *** **** **********
B18hum TCCTCCACCCCTCCTATCTCACCTGCCCAGCCCTGTGCTGGGGCCCTGC B18mouse TCCTCTGACTCTCCC-ATCTCTCCCACCCGGCTCTAGGCC ***** * ***** * **** * * * * * * *
B18hum AATTGGAATGTTGGCCAGATTTCTGCAATAAACACTTGTGGTTTGCGGAA B18mouseAAAAGGCTCGGTACCTCCTTACTGGGAGC-CACGT

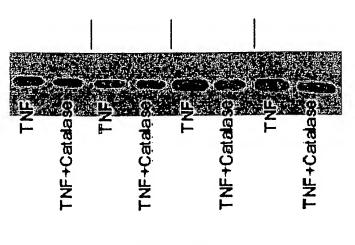
FIG. 5C

## PEROXISOME-ASSOCIATED POLYPEPTIDE...

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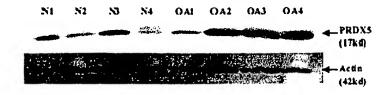


#### PEROXISOME-ASSOCIATED POLYPEPTIDE... Bernard Knoops, et al.

Appl. No.: Unknown Atty Docket: DECLE 30.001 CP1

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FIG. 6A



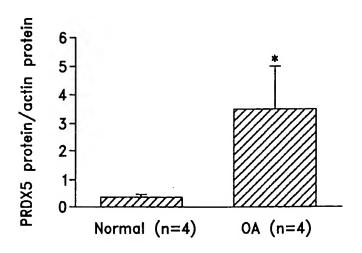
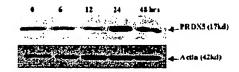
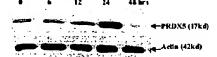


FIG. 6B

FIG. 7A FIG. 7B





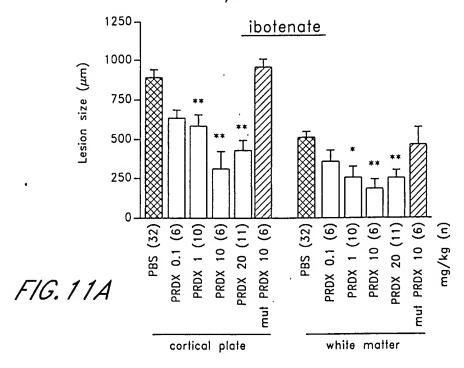
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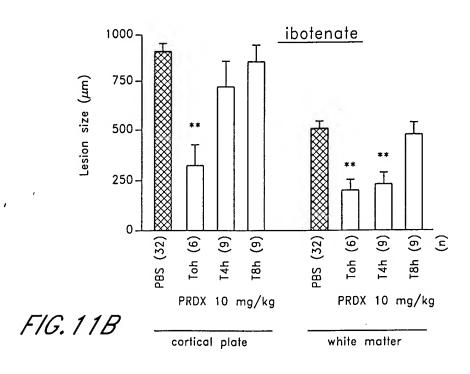


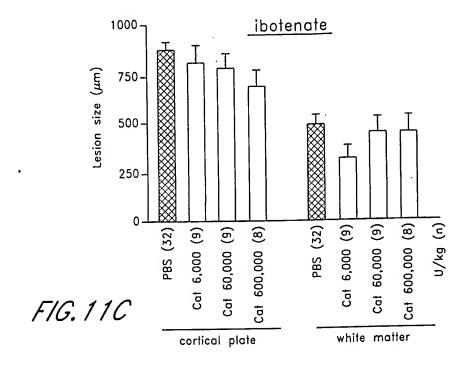
CL

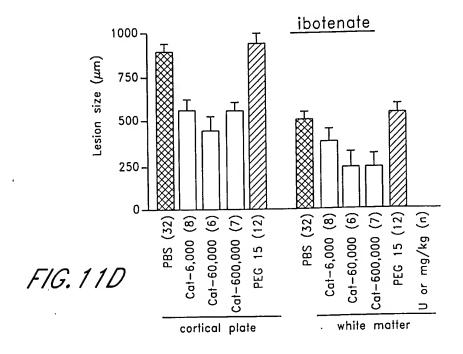
LV

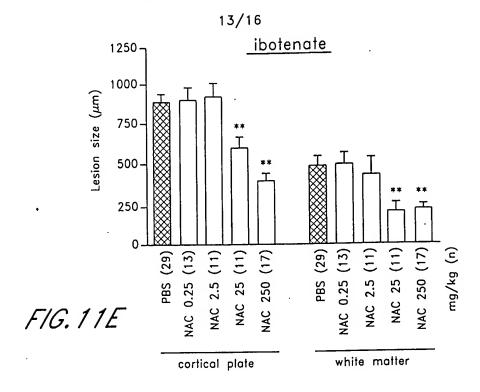
FIG. 10B

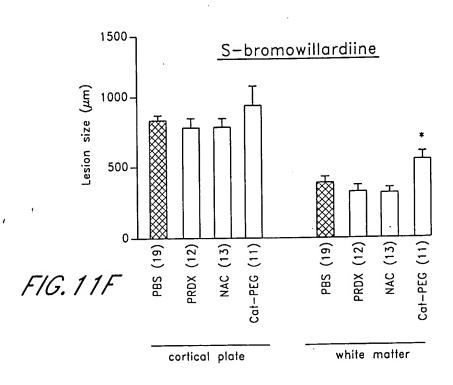


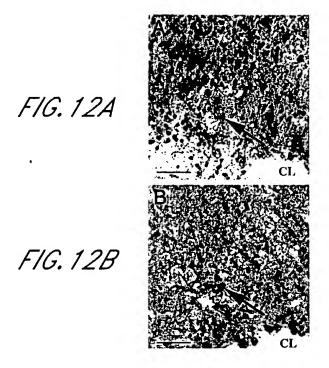












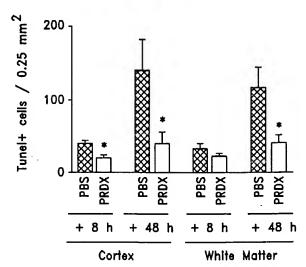


FIG. 12C

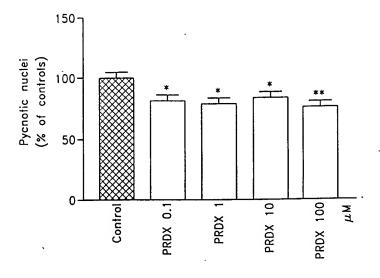
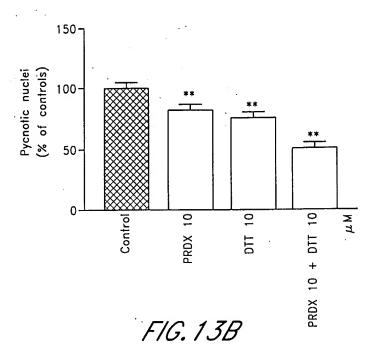


FIG. 13A



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FIG. 1442

FIG. 14B2

A-1
B-1

FIG. 144 1

FIG. 14B 1

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